

Bastian Daubert Presentation Mock-Up

VIAGRA DAUBERT HEARING

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University of California at San Francisco

Prepared: October 3, 2019

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CURRICULUM VITAE**

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1987 - 1987	Dermatologische Klinik, LMU, Munich, Germany	Intern	Dermatology	
1987 - 1988	Schwabinger Krankenhaus, Munich, Germany	Intern	Medicine	
1988 - 1989	Ludwig-Maximilian-University	Fellow	Hematology	
1989 - 1994	University of Würzburg	Resident	Dermatology	
1995 - 1996	University of California, San Francisco	Visiting Scholar	Dermatopathology	
1997 - 1999	University of California, San Francisco; Comprehensive Cancer Center	Fellow	Cancer Genetics Program	

LICENSES, CERTIFICATION

1989 Medical License: Bavarian Medical Board

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- Distinguished Professor of Cancer Biology, Helen Diller Family Comprehensive Cancer Center
- Principal Investigator, Bastian Research Laboratory
- Leader, Cutaneous Oncology Research and Clinical Program
- Director, Molecular Dermatopathology Clinical Service
- Founder and Executive Director, Clinical Cancer Genomics Lab

Boris Bastian, M.D., Ph.D.

Selected Publications on Melanoma Initiation and Progression

The screenshot shows a journal article from The New England Journal of Medicine. The header includes 'REVIEWS' and 'ORIGINAL ARTICLE'. The title of the article is 'Distinct Sets of Genetic Alterations in Melanoma'. The authors listed are John A. Curtin, Ph.D., Jane Fridlyand, Ph.D., Toshiro Kageshita, M.D., Hetal N. Patel, M.S., Klaus J. Busam, M.D., Heinz Kutzner, M.D., Kwang-Hyun Cho, M.D., Setsuya Aiba, M.D., Ph.D., Eva-Bettina Bröcker, M.D., Philip E. LeBoit, M.D., Dan Pinkel, Ph.D., and Boris C. Bastian, M.D. The journal logo and volume information are visible on the left side.

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The NEW ENGLAND JOURNAL OF MEDICINE

ORIGINAL ARTICLE

Distinct Sets of Genetic Alterations in Melanoma

John A. Curtin, Ph.D., Jane Fridlyand, Ph.D., Toshiro Kageshita, M.D., Hetal N. Patel, M.S., Klaus J. Busam, M.D., Heinz Kutzner, M.D., Kwang-Hyun Cho, M.D., Setsuya Aiba, M.D., Ph.D., Eva-Bettina Bröcker, M.D., Philip E. LeBoit, M.D., Dan Pinkel, Ph.D., and Boris C. Bastian, M.D.

ABSTRACT

BACKGROUND
Exposure to ultraviolet light is a major causal relationship between risk and exposure; its heterogeneity is explained by genetically distinct damage and 40 melanomas from skin without sun exposure, and subungual (acral) sites; and 20 mucosal sites.

METHODS
We compared genome-wide alterations in the status of BRAF and N-RAS in 126 melanomas from exposure to ultraviolet light differs: 30 melanomas and 40 melanomas from skin without sun exposure, and subungual (acral) sites; and 20 mucosal sites.

RESULTS
We found significant differences in the frequency of copies of DNA and mutation frequencies in melanomas. Samples could be correctly classified into three groups on the basis of the changes in the number of copies of DNA and mutation frequencies in melanomas. Melanomas arising in skin without sun exposure, and subungual (acral) sites; and 20 mucosal sites, which could be correctly classified into three groups. Eighty-one percent of melanomas from skin with mutations in BRAF or N-RAS, the majority of mutations in neither gene. Melanomas with increases in the number of copies of the genes for cyclin D1 (CDKN1A), downstream components of

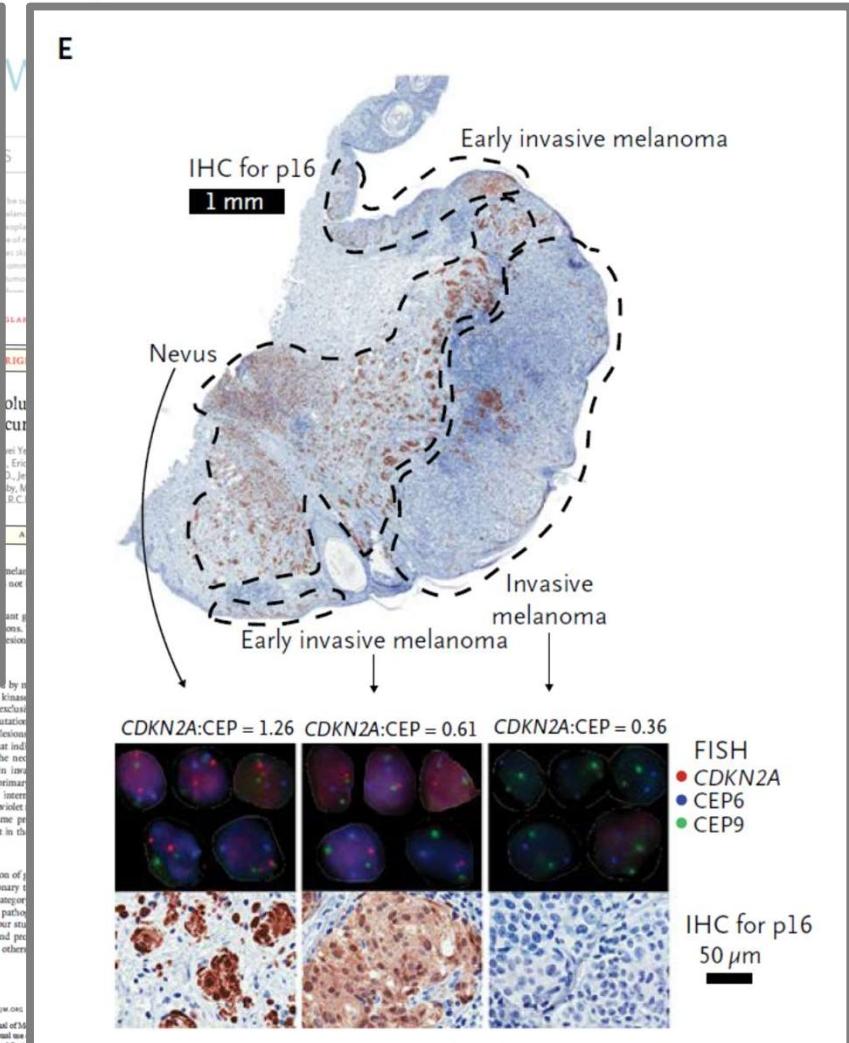
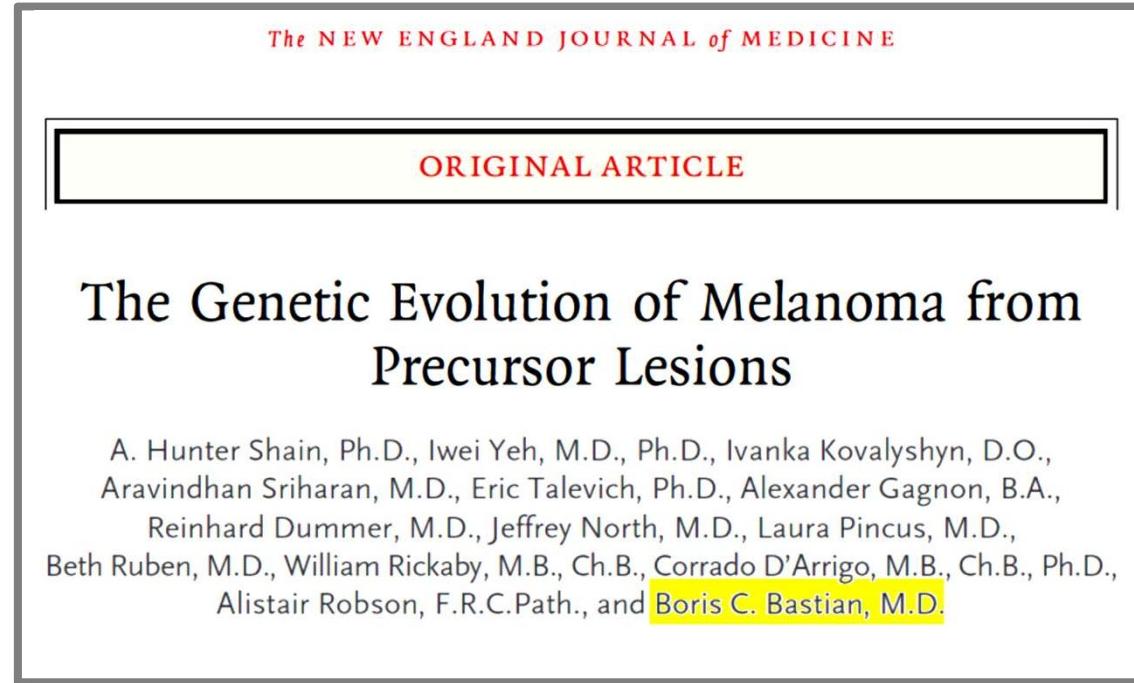
CONCLUSIONS
The genetic alterations identified in melanomas of sun exposure indicate that there are distinct sets of melanomas that implicate CDKN1A and CDKN1B as without mutations in BRAF or N-RAS.

N ENGL J MED 355:20 www.nejm.org

The New England Journal of Medicine
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Boris Bastian, M.D., Ph.D.

Selected Publications on Melanoma Initiation and Progression



Boris Bastian, M.D., Ph.D.

Selected Publications on Melanoma Initiation and Progression



Annu. Rev. Pathol. Mech. Dis. 2014;9:239-271 Downloaded from www.annualreviews.org
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The Molecular Pathology of Melanoma: An Integrated Taxonomy of Melanocytic Neoplasia

Boris C. Bastian

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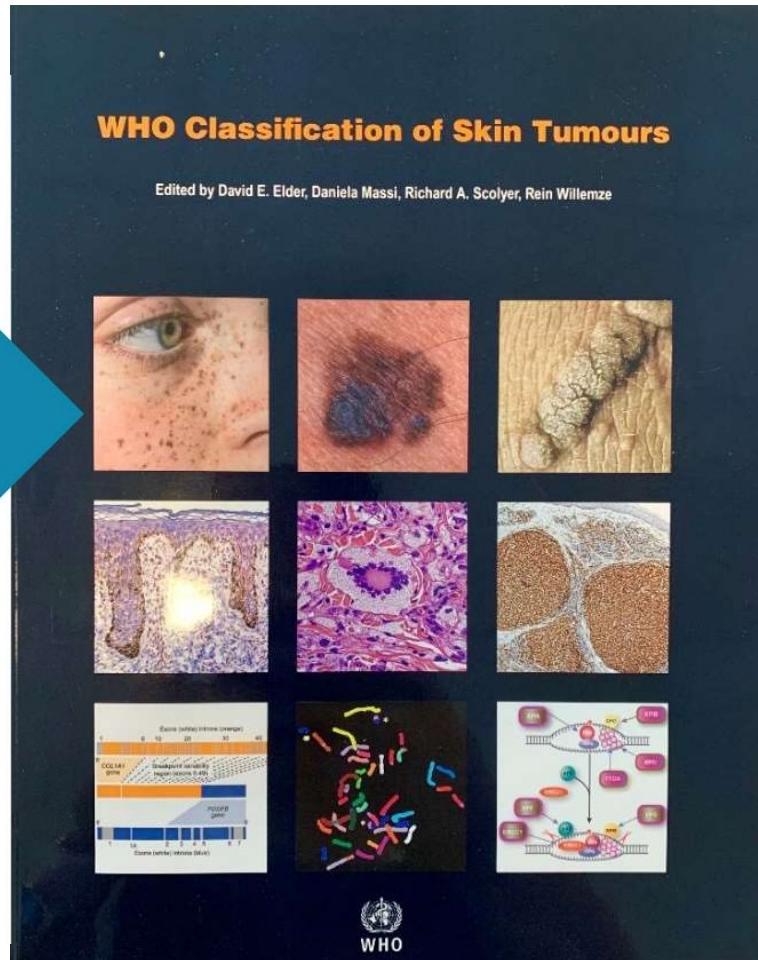
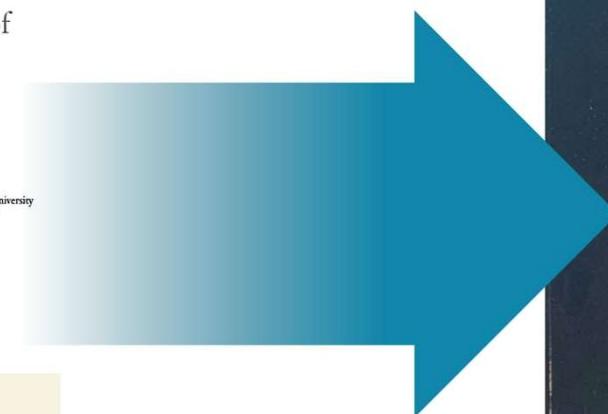
Keywords

genetics, pathogenesis, classification, mutation, nevi

Abstract

Melanomas comprise multiple biologically distinct categories, which differ in cell of origin, age of onset, clinical and histologic presentation, pattern of metastasis, ethnic distribution, causative role of UV radiation, predisposing germ-line alterations, mutational processes, and patterns of somatic mutations. Neoplasms are initiated by gain-of-function mutations in one of several primary oncogenes, which typically lead to benign melanocytic nevi with characteristic histologic features. The progression of nevi is restrained by multiple tumor-suppressive mechanisms. Secondary genetic alterations override these barriers and promote intermediate or overtly malignant tumors along distinct progression trajectories. The current knowledge about the pathogenesis and clinical, histologic, and genetic features of primary melanocytic neoplasms is reviewed and integrated into a taxonomic framework.

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Boris Bastian, M.D., Ph.D.

Selected Publications on Melanoma Initiation and Progression

Cancer Cell

Bi-allelic Loss of CDKN2A Initiates Melanoma Invasion via BRN2 Activation

Graphical Abstract

Authors
Hanlin Zeng, Aparna Jorapur, A. Hunter Shain, ..., Iwei Yeh, Boris C. Bastian, Robert L. Judson

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In Brief
Zeng et al. find that complete *CDKN2A* loss coincides with the onset of invasiveness in melanocytic tumors at distinct progression stages. *p16^{INK4A}*, encoded by *CDKN2A*, inhibits E2F1-mediated transcriptional activation of *BRN2*, a transcription factor that has been associated with melanocytic invasive programs.

Background
Exposure to sunlight is the major risk factor for melanoma. We compared the genomic status of *BRN2* in melanomas from patients with and without a history of sun exposure to identify mutations in *BRN2* that are associated with increased invasiveness. We found significant increases in the number of copies of *BRN2* in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure. We found that the increase in *BRN2* copy number was associated with increased invasiveness in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure.

Methods
We compared the genomic status of *BRN2* in melanomas from patients with and without a history of sun exposure to identify mutations in *BRN2* that are associated with increased invasiveness. We found significant increases in the number of copies of *BRN2* in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure.

Results
We found significant increases in the number of copies of *BRN2* in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure. We found significant increases in the number of copies of *BRN2* in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure.

Conclusion
The genetic analysis of melanomas from patients with and without a history of sun exposure revealed that the increase in *BRN2* copy number is associated with increased invasiveness in melanomas from patients with a history of sun exposure compared with those without a history of sun exposure.

Cancer Cell

Genomic and Transcriptomic Analysis Reveals Incremental Disruption of Key Signaling Pathways during Melanoma Evolution

Graphical Abstract

Article

Genomic and Transcriptomic Analysis Reveals Incremental Disruption of Key Signaling Pathways during Melanoma Evolution

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Cancer Cell

Bi-allelic Loss of CDKN2A Initiates Melanoma Invasion via BRN2 Activation

Graphical Abstract

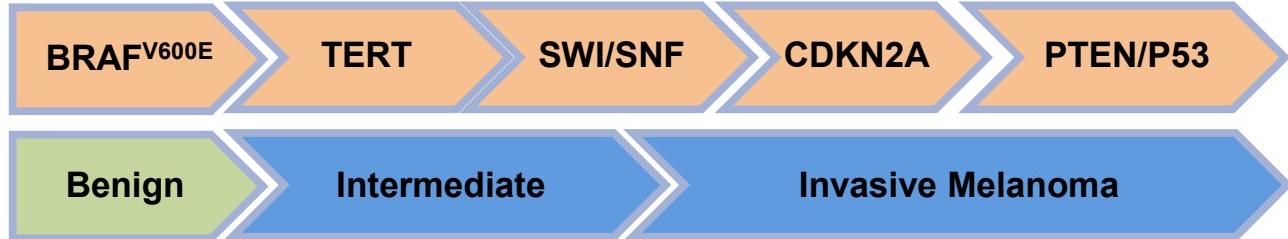
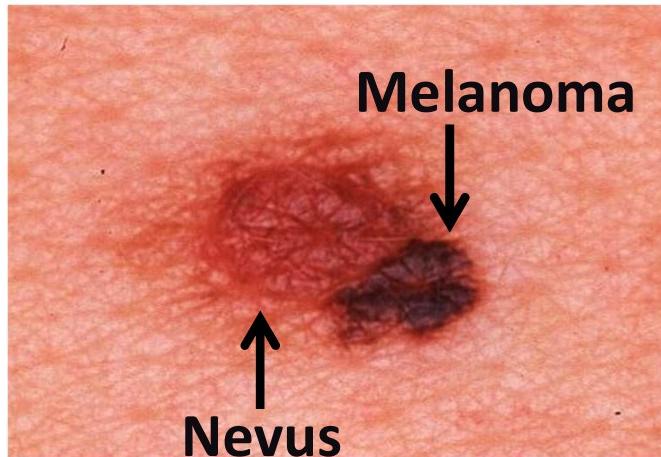
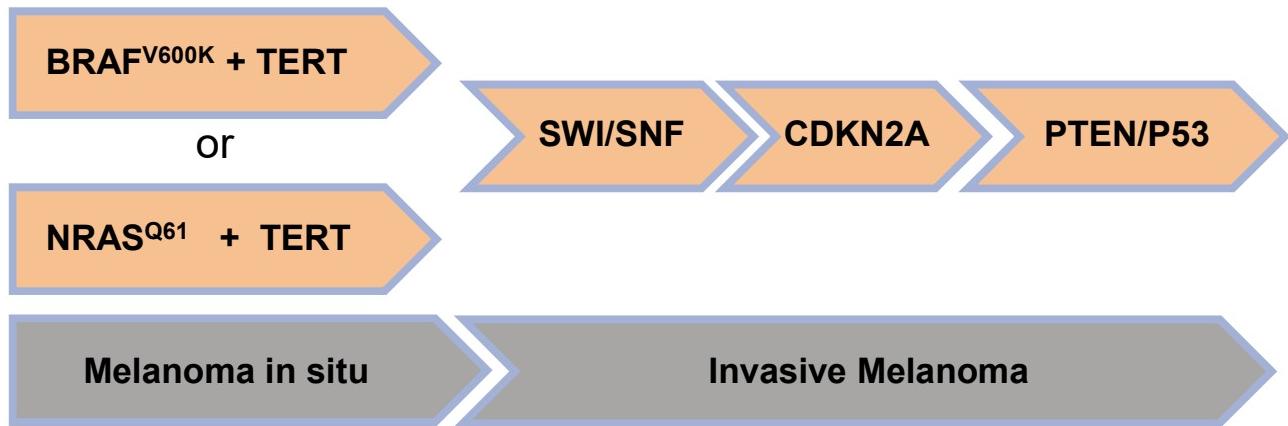
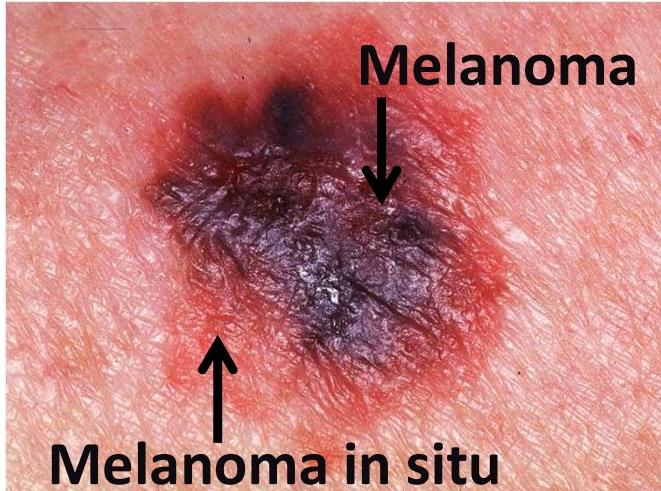
Significance
Engineering of human melanocytes is a tractable model for melanoma initiation.

- The *CDKN2A* locus suppresses melanocyte migration and melanoma invasion.
- p16^{INK4A}* loss drives melanoma invasion via *BRN2* activation.
- BRN2* is a direct transcriptional target of E2F1.

CellPress

Zeng et al., 2018, *Cancer Cell* 34, 56–68
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<https://doi.org/10.1016/j.ccr.2018.05.014>

A Stepwise Accumulation of Mutations Causes Melanoma Initiation and Progression



“No Causal Link between Phosphodiesterase Type 5 Inhibition and Melanoma”

No Causal Link between Phosphodies Inhibition and Melanoma

Jenny Z. Wang¹, Stephanie Le², Claire Alexanian¹, Sucharita Boddu¹, Alia Alina Marusina², Emanuel Maverakis¹

¹Albert Einstein College of Medicine, Bronx, NY; ²Department of Dermatology, University of California CA; ³Georgetown University School of Medicine, Washington, DC, USA

Purpose: To examine the association between phosphodiesterase type 5 (PDE5) inhibitor use and a systematic review of observational studies; and 2) determining if low PDE5A gene expression related with decreased overall survival.

Materials and Methods: A systematic search of observational studies examining the association between PDE5 inhibitor use and melanoma was performed through ClinicalTrials.gov, the Cochrane Library, EMBASE, PubMed, and seven eligible studies were identified. PDE5A gene expression was analyzed with human melanoma samples obtained from The Cancer Genome Atlas.

Results: Four studies reported a positive association between PDE5 inhibitor use and melanoma. RNA sequencing data analysis revealed that under-expression of the PDE5A gene comes in melanoma.

Conclusions: There is currently no evidence to suggest that PDE5 inhibition in patients causes melanoma. The few observational studies that demonstrated a positive association between PDE5 failed to account for major confounders. Nonetheless, the substantial evidence implicating PDE5 inhibitors in the treatment of penile erectile dysfunction warrants further investigation.

Keywords: Melanoma; Phosphodiesterase 5 inhibitors; Sildenafil citrate; Tadalafil; Vardenafil dihydrochloride

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INTRODUCTION

Ever since the introduction of sildenafil 20 years ago, phosphodiesterase type 5 (PDE5) inhibitors have become the mainstay therapy in the treatment of penile erectile dysfunction (ED), which can be severely limiting in an estimated 5% to 20% of men worldwide [1].

PDE5 inhibitors work by an activity that degrades cGMP, a smooth muscle caverousum of the penis, resulting in penile smooth muscle relaxation enhancement of the link between PDE5 and ED.

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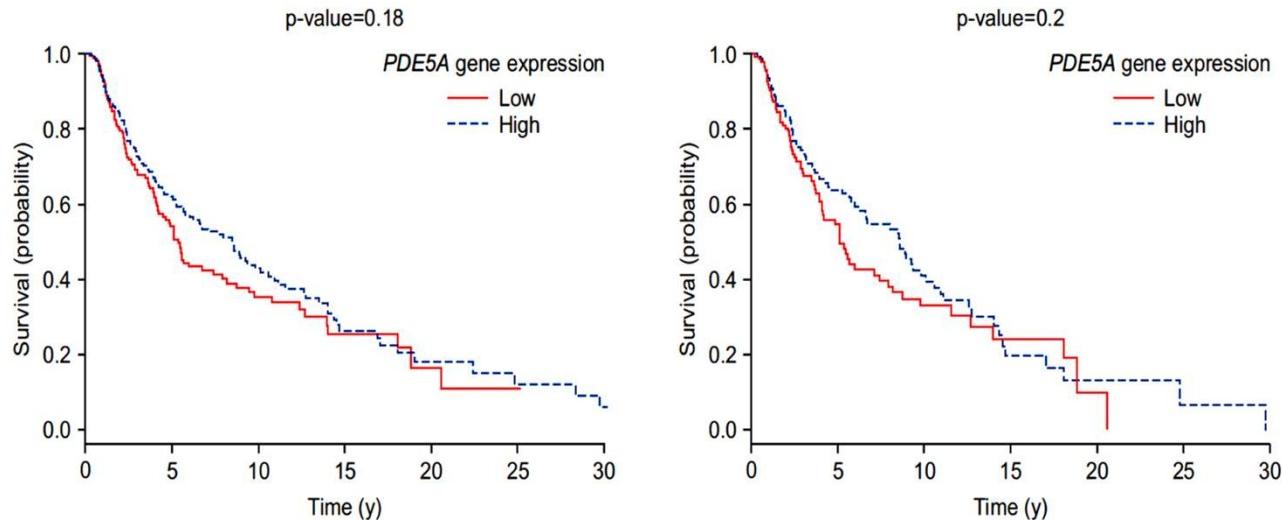


Fig. 1. Kaplan-Meier survival curve for differential expression levels of PDE5A gene in (A) 470 patients (180 females and 290 males) with a diagnosis of melanoma at any tumor stage (0–IV), ages 14–91 years; and (B) specifically male melanoma patients (n=287) with a diagnosis of melanoma at any tumor stage (0–IV), ages 18–91 years. Melanoma prognosis unaffected by high or low PDE5A expression in patients, regardless of gender, age, or tumor stage.

Arozarena *In Vivo*: No Invasion Effect

Case 3:16-md-02691-RS Document 837-35 Filed 01/11/19 Page 1 of 21

Cancer Cell
Article

Oncogenic BRAF Induces Melanoma Cell Invasion by Downregulating the cGMP-Specific Phosphodiesterase PDE5A

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DOI 10.1016/j.ccr.2010.09.029

SUMMARY

We show that in melanoma cells oncogenic BRAF, which downregulates the cGMP-specific phosphodiesterase, has a small decrease in proliferation, its major impact is to stimulate invasion. This is because PDE5A downregulation leads to an increase in Ca^{2+} levels, stimulating increased contractility and inducing an increase in short-term and long-term colonization of the pathway in NRAS mutant melanoma or BRAF mutant cancer cells. Oncogenic BRAF induces invasion through downregulation of PDE5A.

INTRODUCTION

Melanocytes are specialized pigment cells located primarily in the skin, where they determine complexion and hair color and provide protection from the damaging effects of ultraviolet radiation. Melanocytes are also found in the eye and gut. These cells are also the precursors of melanoma, a potentially deadly skin cancer that kills about 8,000 people in the United States and about 12,000 people in Europe each year. In many Western societies, melanoma incidence almost doubles every decade. If treated early, melanoma can be cured by surgical resection, but due to its proclivity to metastasize, in about 20% of patients it progresses to an aggressive invasive disease that is refractory to treatment and has a poor prognosis, with median survival rates of 6–9 months and 5 year survival rates of 5%–10%. These data highlight the need for improved understanding of melanoma biology to facilitate development of therapeutic strategies.

An important signaling pathway in melanoma is the RAS/RAF/MEK/ERK cascade (Gray-Shocher et al., 2007). RAS is a small G

Significance

The protein kinase BRAF is activated by somatic gain-of-function mutations in ERK pathway hyper-activation, and we show that this oncogene activates PDE5A. PDE5A is the target of drugs such as sildenafil, tadalafil, and vardenafil, and is involved in the pathophysiology of pulmonary arterial hypertension. PDE5A downregulation results in increased cell invasion both *in vitro* and *in vivo*, and increased long-term colonization of melanoma cells. Our findings suggest that PDE5A may be a key target for anti-metastatic therapy.

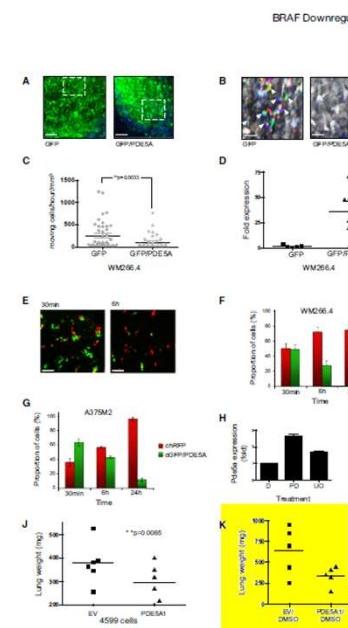


Figure 7. PDE5A Regulates Melanoma Cell Invasion In Vivo
 (A) Low-resolution still images taken from video recordings of subcutaneous cultures formed from WM266.4-GFP/PDE5A or WM266.4-GFP/PDE5A-GFP/PDE5A cells. Scale bar, 100 μm . (B) High-resolution images of the cells shown in the dotted boxes in (A). The scale bar of the cells were taken at 10, 1.5, and 20 μm . (C) Quantification of moving cells per frame in the field of view. (D) Quantification of moving cells per frame from 3 h cultures formed using WM266.4-GFP or WM266.4-GFP/PDE5A (GFP/PDE5A) cells. The solid bars represent the average number of moving cells for the two populations with error bars to represent standard deviations from the mean. (E) PDE5A mRNA expression in WM266.4-GFP (GFP) or WM266.4-GFP/PDE5A (GFP/PDE5A) cells was determined by RT-PCR. Five times for each genotype (triplicate, and averaged) and individual tumors are shown. Relative to the value of the PDE5A1/GFP control. The bars represent the average level of relative gene expression. (F) Quantification of WM266.4-GFP cells in WM266.4-GFP/PDE5A (green) cells in the lungs of mice 30 min or 6 hr after injection with equal numbers of each line. Scale bar, 75 μm . (G) Quantification of 10 fields of cells from 3 mice 30 min, 6 hr, or 24 hr after injection with equal numbers of WM266.4-dsRIP (dsRIP) or WM266.4-GFP/PDE5A (GFP/PDE5A) cells. (H) Quantification of 10 fields of cells from 3 mice 30 min, 6 hr, or 24 hr after injection with equal numbers of A3789 cells (GFP) or A3789-GFP/PDE5A (GFP/PDE5A) cells. (I) Data from the same experiment as (H) except that Pde5a mRNA was inhibited using 4599 mouse melanoma cells treated with DMSO (D), PDE5A1 (P), or U0126 (U, 10 μM) for 24 hr. (J) Western blot showing PDE5A1 and ERK (loading control) levels in 4599 melanoma cells engineered for stable expression of PDE5A1 or an empty vector (EV) control. (K) Lung weights from mice following tail vein injection of 4599 melanoma cells expressing empty vector (EV) or PDE5A1. The weights of the individual lungs are shown, with the bars representing the mean

54 Cancer Cell 19, 45–57, January 18, 2011 ©2011 Elsevier Inc.

Research Community on B16 Mouse Melanoma



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Pigment Cell Melanoma Res. 2013 July; 26(4): E8–E14. doi:10.1111/pcmr.12088

The Future of Preclinical Mouse Models Is Now

Glenn Merlino, Keith Flaherty, Nicolas Acciari, Terry Van Dyke, and Meenhard Herlyn

On October 17th, 2012 a rela scientists met at the Wistar I the current status and future; on the mouse as a model sys (NCI), Terry Van Dyk (NC development of better precl tell modelers what informati modelers acquaint clinicians scientists included Drs. Keit (Memorial Sloan Kettering), and Nicolas Acuña-Pes researchers included Drs. Ar Sean Holmen (Huntsman C Keiran Smalley (Moffitt Ca (CSO Adelton Medical Rese Foundation Breakthrough Ce Research Alliance).

An underlying premise of this without a solid rationale denials always seem to be one support them. Candidate and that typically employ the graft subcutaneously xenografted models are overly reliant on culture and have an inadequate system, and so have proven many drugs move into the clinical melanoma field finds its predictive of clinical response discussions on the current set models, and how these two descriptions of new resources and conclusions concerning challenges.

However, current in depth knowledge about the complex molecular basis of human melanoma has revealed that this well-established and widely used cell line [B16] **does not reflect the genetic underpinnings of human melanoma.**

the cognate immune system for any model may be paramount in developing accurate immunological response hypotheses.

For more than five decades, the tumor immunology field has greatly benefited from the development of the poorly immunogenic B16 mouse melanoma model. However, current in-depth knowledge about the complex molecular basis of human melanoma has revealed that this well-established and widely used cell line does not reflect the genetic underpinnings of human melanoma. Instead, preclinical animal models used for efficacy testing of novel immunotherapeutic modalities should be founded on relevant human biology. To help achieve this, the malignant landscape of human melanoma should be modeled on mouse hosts with intact immune systems (e.g., C57BL/6 mice), allowing the evaluation of the effects of the immune system and host-tumor interactions. Ideally, a tumor exhibiting intrinsic immunogenicity on an immunocompetent host would result in the intratumoral recruitment of cellular elements comprising both the innate and adaptive arms of the host immune system. This desired attribute could recapitulate the failure of host immunologic barriers to impede effective antitumor responses in humans.

In addition, the presence of tumor infiltrating lymphocytes in immunocompetent mouse models would recapitulate the occurrence of functionally tolerant T-cell repertoires against unknown tumor antigens in patients with melanoma. A point of concern is the insertion of artificial genetic information (e.g. "non-self" proteins) into tumors that may inadvertently increase their immunogenicity. The artificial aspects of this sort of immunity may not

Dhayade *In Vitro*: Does Not Study Sildenafil Alone

Case 3:16-md-02691-RS Document 837-37 Filed 01/11/19 Page 1 of 26

Cell Reports

Sildenafil Potentiates a cGMP-Dependent Pathway to Promote Melanoma Growth

Graphical Abstract

Authors

Case 3:16-md

Highlights

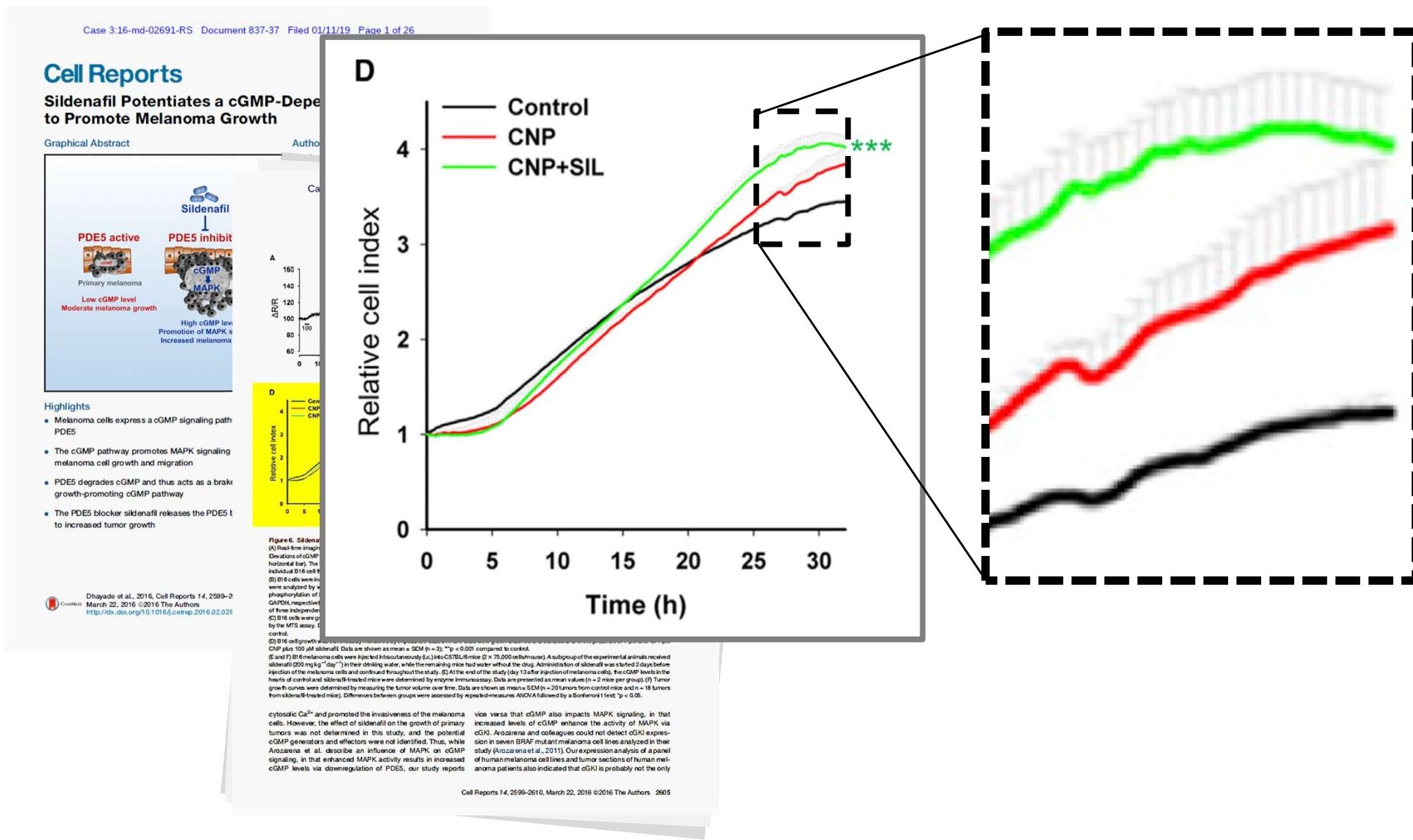
- Melanoma cells express a cGMP signaling pathway PDE5
- The cGMP pathway promotes MAPK signaling in melanoma cell growth and migration
- PDE5 degrades cGMP and thus acts as a brake growth-promoting cGMP pathway
- The PDE5 blocker sildenafil releases the PDE5 to increase tumor growth

Figure S5 (Related to Figure 7). Model of cGMP Signaling in Melanoma Pathogenesis

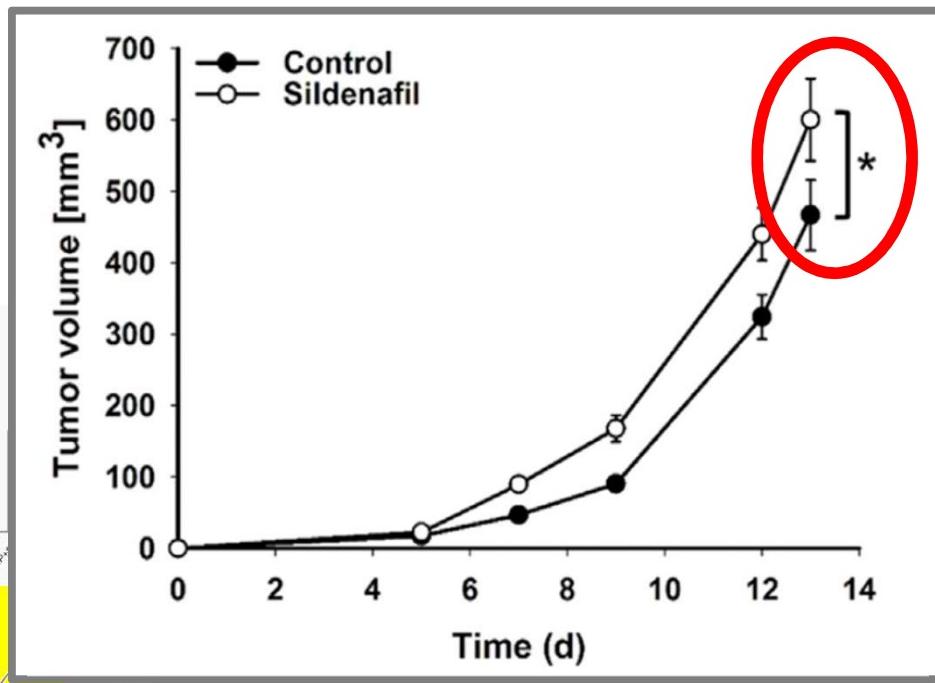
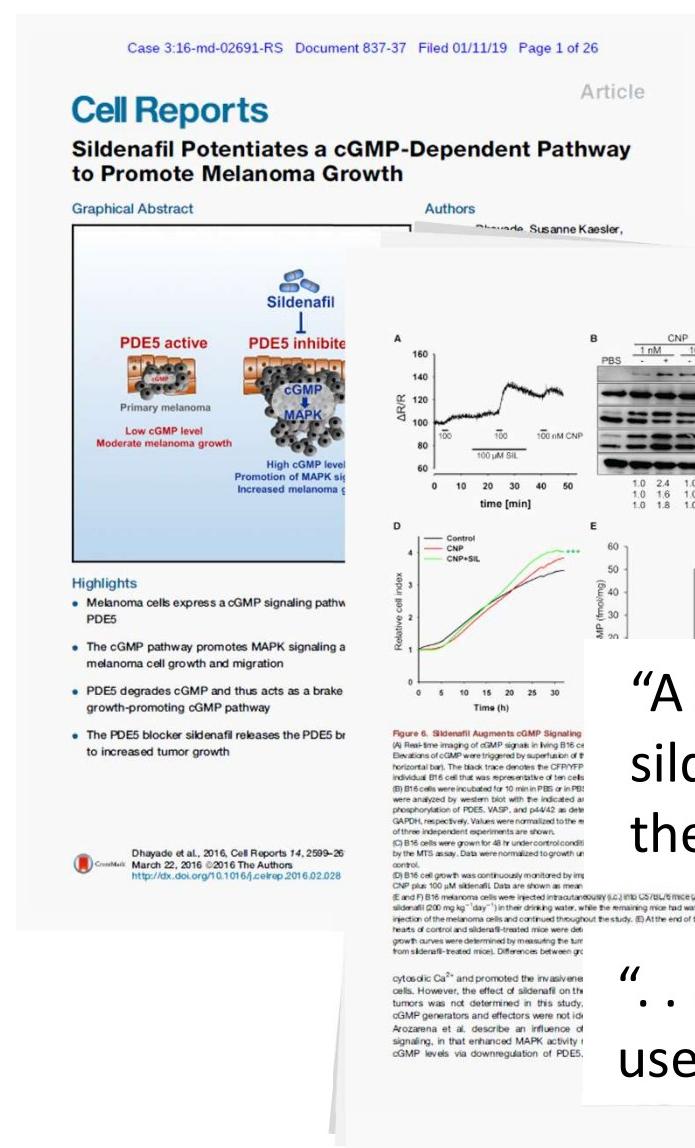
The scheme illustrates how a bidirectional crosstalk of cGMP and MAPK signaling promotes the switch from non-metastatic cells in primary melanomas to invasive/metastatic cells. CNP is released from endothelial cells (dashed circle) of the inflamed tumor vasculature (upper left) and binds to its receptor, GC-B, on melanoma cells. Thereby, CNP triggers an increase of the intracellular cGMP concentration and activation of cGKII in the melanoma cells. Via phosphorylation of yet unknown substrate proteins, cGKII promotes MAPK signaling upstream of MEK (indicated by the bracket), resulting in cells with increased potential for growth, migration, and invasiveness to develop. Degradation of cGMP via PDE5 acts as a "break" in this switching process. However, a persistent increase in MAPK signaling, for instance, by sustained activity of the CNP-cGMP-cGKII cascade and/or by somatic gain-of-function mutation of BRAF to V600E-BRAF (Arozarena et al., 2011), results in downregulation of PDE5 at the transcriptional level. This releases the PDE5 "break", thus, establishing a feed-forward, self-reinforcing loop that further enhances the aggressiveness of the melanoma cells. The PDE5 "break" can also be released pharmacologically by the PDE5 inhibitor sildenafil. Note that CNP and sildenafil might act mainly on cells of the primary tumor to increase their metastatic potential, and that "switched" metastatic cells might lose expression of both cGKI and PDE5. Note also that the present study has analyzed the effects of exogenously supplied CNP. The role of inflamed tumor vessels as the source of endogenous CNP in melanomas needs to be established in future studies. U0126 is a MEK inhibitor used in the present study. RTK stands for receptor tyrosine kinase.

The role of inflamed tumor vessels as the **source of endogenous CNP** in melanomas needs to be **established in future studies.**

Dhayade *In Vitro*: Does Not Study Sildenafil Alone



Supratherapeutic Sildenafil Dosing



“A subgroup of the experimental animals received sildenafil (**200 mg kg/day**) in their drinking water, while the remaining mice had water without the drug.”

“... it is **not clear** whether the sildenafil concentration used in our experiments is also reached **in patients**. . .”

Direct Effects: PDE-5 Inhibitors and Melanoma

ARTICLE

Journal of Cellular Biochemistry 113:2738–2743 (2012)

Journal of Cellular
Biochem[®]

PDE5 Inhibitor Promotes Melanin Synthesis Through PKG Pathway in B16 Melanoma Cells

Xiaodong Zhang,¹ Guirui Yan,¹ Jun Ji,^{1,2} Jingwei Wu,¹ Xiaoyun Sun,¹ Jingshan Shen,¹ Huaoliang Jiang,¹ and Heyao Wang^{1*}

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ABSTRACT

PDE inhibitors could increase cellular cGMP levels and are used to treat erectile dysfunction as well as pulmonary arterial hypertension. It was reported that PDE5 inhibitor is necessary for UVB-induced melanin synthesis, however, the effect of PDE5 inhibitor on melanin synthesis has not been examined. We found that PDE5 inhibitor (sildenafil or vardenafil) and the cGMP analog 8-CPT-cGMP phosphorylation, leading to increased tyrosinase expression and melanin synthesis, which was counteracted by KT5823 (cGMP-dependent protein kinase (PKG) inhibitor). However, KT5823 did not affect cAMP-elevating agent-mediated melanin synthesis. Our results suggest that PDE5 inhibitor may be beneficial for the treatment of hypopigmentation diseases. *J. Cell. Biochem.* 113: 2738–2743, 2012. © 2012 Wiley Periodicals, Inc.

KEY WORDS: MELANIN SYNTHESIS; PDES INHIBITOR; PKG; TYROSINASE

Melanin is an important skin pigment in human and contributes significantly to the health of an individual [Gilkrest, 1989; Sturm, 2002]. Lack or decreased levels of melanin in human lead to many skin diseases, termed hypopigmentary disorders including vitiligo and gray hair [Hartmann et al., 2004; Dessimoni et al., 2009]. There are several treatments for hypopigmentary diseases, but there remain some problems, such as poor efficacy and severe side effects [Hartmann et al., 2004; Hercogova et al., 2007]. Thus the search for new types of treatments of hypopigmentary diseases with high efficacy and low toxicity is warranted.

Melanin is synthesized in melanocytes via a cascade of enzymatic reactions controlled by tyrosinase [Hearing, 1999; Kim et al., 2010]. Tyrosinase is the rate-limiting enzyme of melanin synthesis that catalyzes the hydroxylation of L-tyrosine to L-3,4-dihydroxyphenylalanine (L-DOPA) and the oxidation of L-DOPA to dopaquinone

[Slominski et al., 2004]. Melanin synthesis is still variable of intrinsic and extrinsic factors, including agents, UVB, and a wide variety of growth factor [Friedmann and Gilkrest, 1987; Yasumoto et al., 2009; Hu et al., 2010]. cAMP pathway plays a regulation of melanogenesis through activation of protein kinase (PKA) and cAMP response element transcription factor, which induced an up-regulation expression and the stimulation of melanogenesis et al., 2004; Park et al., 2009].

For the first time in 1996, the second messenger to be required for melanin synthesis induced by increase cGMP content in melanocytes [Rome 1996]. Phosphodiesterase 5 (PDE5) is the responsible for cGMP hydrolysis in various type [Kass et al., 2007]. PDE5 inhibitors such as silder-

Xiaodong Zhang and Guirui Yan contributed equally to this work.

Additional supporting information may be found in the online version of this article.

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3',5'-Cyclic Nucleotide Phosphodiesterase in Tumor Cells as Potential Target for Tumor Growth Inhibition

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ABSTRACT

Isoenzymes of 3',5'-cyclic nucleotide phosphodiesterase (PDE) have been characterized in B16 murine melanoma cells and MCF-7 human mammary carcinoma cells. Separation of soluble phosphodiesterase activity by fast protein liquid chromatography on a Mono-Q column resolved three isoenzymes, MCF-7 cells contained a cyclic GMP-specific isoenzyme (PDE-V), a cyclic GMP-activatable isoenzyme (PDE-II), and a cyclic AMP-specific isoenzyme (PDE-IV). B16 cells contained a cyclic GMP-specific isoenzyme (PDE-V), a Ca²⁺/calmodulin-activated isoenzyme (PDE-I), and a cyclic AMP-specific isoenzyme (PDE-IV).

A series of PDE inhibitors was tested for their activity spectrum on PDE isoenzymes. Inhibition of PDE activity in B16 cells by the new compound DC-TA-46, was found to result specifically from PDE-IV inhibition (50% inhibition (IC_{50}) = 0.03 μ M). Much lower inhibitory activity was observed for DC-TA-46 toward PDE-I (IC_{50} = 5 μ M) and PDE-V (IC_{50} = 14 μ M).

DC-TA-46 was found to inhibit growth of B16 melanoma and MCF-7 mammary carcinoma cells dose-dependently (B16: IC_{50} = 1.7 μ M, MCF-7: IC_{50} = 2 μ M). At 2 μ M concentration, growth inhibition of B16 melanoma cells was 60%, concomitant with a decrease in PDE activity of 63% and an increase in cAMP level of 59%. In contrast, incubation with inhibitors specific for PDE-I and PDE-V resulted only in marginal or undetectable growth inhibition. The results suggest a correlation between PDE-IV inhibition and growth inhibition. PDE-IV thus appears to be a potential new target for antiproliferative treatment.

INTRODUCTION

Multiple forms of PDE¹ have been demonstrated in various tissues or cells and have been characterized on the basis of substrate specificity, sensitivity to calmodulin or phosphodiesterase inhibitors, and kinetic parameters (1–3). However, no data are available yet concerning isoenzyme distribution in MCF-7 and B16 tumor cells. We found three isoenzymes: MCF-7 cells, PDE-II, PDE-IV, and PDE-V; in B16 cells, PDE-I, PDE-IV, and PDE-V. Isoenzyme nomenclature follows that introduced by Beavo [2002].

Each form of isoenzyme has a unique role in the regulation of the intracellular level of cyclic nucleotides. cAMP is a positive intracellular signal for cell proliferation in many differentiated cells (4, 5). In many tumor cells, however, cAMP is a negative messenger for proliferation, showing a much lower basal level than in normal cells (5). Some data indicate that the activity of 3',5'-cyclic nucleotide phosphodiesterase is elevated in tumor cells (3, 6). Various agents elevating cAMP have previously been found to inhibit tumor cell growth *in vitro*. PDE inhibitors, especially those of the methylxanthine type, display, however, growth inhibition only at rather high concentrations (5, 7–14).

A new PDE inhibitor, DC-TA-46 (15), with potent inhibitory activity toward PDE from B16 melanoma and MCF-7 mammary carcinoma

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MATERIALS AND METHODS

Preparation of Cell Extracts. Cells were performed at 0°C buffer, was suspended in 1 hexamidine, 0.1 mM phenylmethylsulfonyl peptide, 0.25 mM sucrose, 0.05 at 1,000 \times g for 10 min. T 100,000 \times g for 60 min to PDE Assay. PDE activity (16) with slight modification in buffer containing 50 mM sucrose was carried out: 0.1 mM ³H-cAMP (IC₅₀ = 0.03 μ M). Much lower inhibitory activity was observed for DC-TA-46 toward PDE-I (IC₅₀ = 5 μ M) and PDE-V (IC₅₀ = 14 μ M).

DC-TA-46 was found to inhibit growth of B16 melanoma and MCF-7 mammary carcinoma cells dose-dependently (B16: IC_{50} = 1.7 μ M, MCF-7: IC_{50} = 2 μ M). At 2 μ M concentration, growth inhibition of B16 melanoma cells was 60%, concomitant with a decrease in PDE activity of 63% and an increase in cAMP level of 59%. In contrast, incubation with inhibitors specific for PDE-I and PDE-V resulted only in marginal or undetectable growth inhibition. The results suggest a correlation between PDE-IV inhibition and growth inhibition. PDE-IV thus appears to be a potential new target for antiproliferative treatment.

Monooxygenase Assay. PDE activity was measured at 1 (1 \times 0.5 cm), preincubated 0.1 mM phenylmethylsulfonyl peptide, 1-(4-chlorophenyl)-4-(chloromethyl)cyclotriphosphazene (Tris/HCl, pH 7.4). After washout at 1 min, using a solution of 1 mM were collected.

Effects of Ca²⁺/Calmodulin. Determination of IC₅₀ was performed in the presence of GMP.

Determination of IC₅₀. Cells were washed at a stock solution B to provide a range of concentration in the assay medium was replaced every 48 h throughout performed in triplicate.

Kinetic Parameters. 1 eluting from the Mono-Q column. Final concentration of 10 nM to 100 of 500 μ M. A that no more than 15% of from Hanes and Lineweaver Cell Culture. Cells CO₂, B16 cells were grown per same medium containing Cells (7.2 \times 10⁵) were replated every 48 h throughout performed in triplicate, and was 100%.

Cell Culture. Cells CO₂, B16 cells were grown per same medium containing Cells (7.2 \times 10⁵) were replated every 48 h throughout performed in triplicate, and was 100%.

ANTICANCER RESEARCH 30: 355–358 (2010)

Expression and Role of Phosphodiesterase 5 in Human Malignant Melanoma Cell Line

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Abstract. **Background:** Eleven phosphodiesterase (PDE) gene families (PDE-I–II) have been identified, and some PDE isoforms are selectively expressed in various cell types. Previously, we reported PDE1, PDE3 and PDE4 expressions in human malignant melanoma cells. However, the expression and role of PDE5 in malignant melanoma cells is not clear. Therefore, we characterized PDE5 in human malignant melanoma MAA cells. Materials and Methods: PDE5 activity and PDE5A mRNA expression were investigated in MAA cells. The full open reading frames for human PDE5A were sequenced. Effects of PDE5 inhibitors on cell growth were determined by 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium, inner salt (MTS) assays. Results: PDE5 activity and PDE5A mRNA expression were detected in MAA cells. The nucleotide sequence of PDE5A was identical to that of human PDE5A, previously published. Two PDE5 inhibitors inhibited the growth of cells. Conclusions: PDE5A mRNA is expressed and may play an important role in the growth of human malignant melanoma MAA cells.

Materials and Methods

Cell culture. Human malignant melanoma MAA cells were established and maintained in RPMI 1640 containing 10% fetal bovine serum (Invitrogen Corp., Carlsbad, CA, USA) at 37°C in a humidified 5% CO₂ atmosphere in our laboratory (6).

cGMP/PDE activity assay in cell extracts. The cells were seeded at 1 \times 10⁶ cells/25-cm² flask. After 3 days, the cells were washed twice with phosphate-buffered saline (PBS), harvested with a rubber policeman, and homogenized in ice-cold homogenization buffer (1 mL) with T230 cell-freezing homogenizer (Tomy Seikagaku, Tokyo, Japan).

the hydrolysis of cyclic nucleotides, PDEs regulate the intracellular concentrations and effects of these secondary messengers. Some PDE families are relatively specific for cAMP (PDEs 4, 7 and 8) or for cGMP (PDEs 5, 6 and 9); others hydrolyze both (PDEs 1–3, 10 and 11) (1, 2, 4).

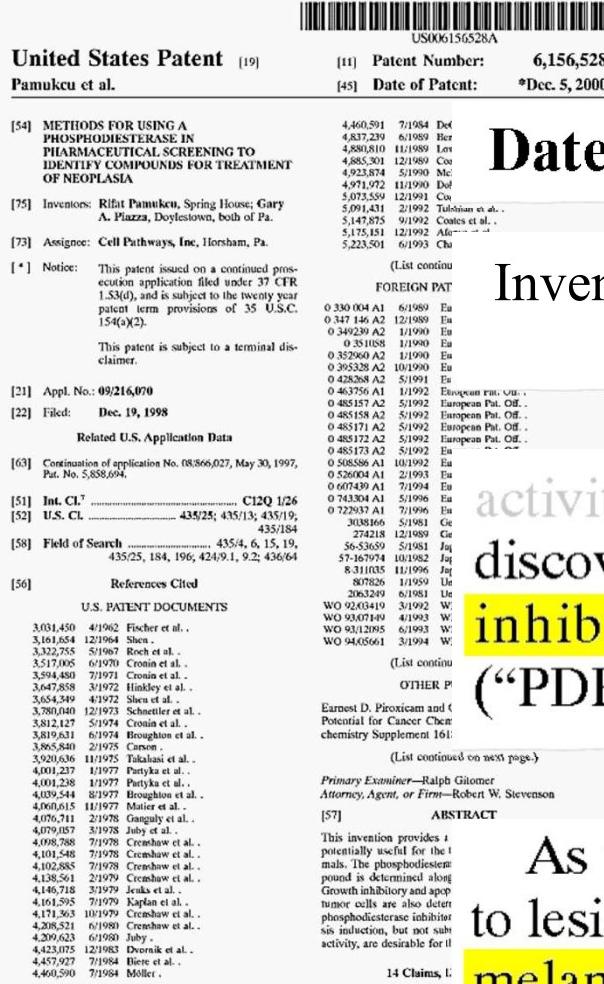
PDE5 is relatively specific for cGMP and is expressed abundantly in vascular smooth muscle, including the pulmonary vasculature and corpus cavernosum of the penis. Three alternatively splicing variants of human PDE5A, PDE5A2, and PDE5A3 have been identified and their tissue distribution differs (2, 3, 5). PDE5 inhibitor sildenafil improves penile erection with a minimal risk of side-effects and adverse events in many men with erectile dysfunction (1–3, 5). However, the expression and role of PDE5 in human malignant melanoma cells is not clear. Therefore, we examined PDE5 in human malignant melanoma MAA cells.

Materials and Methods

Cell culture. Human malignant melanoma MAA cells were established and maintained in RPMI 1640 containing 10% fetal bovine serum (Invitrogen Corp., Carlsbad, CA, USA) at 37°C in a humidified 5% CO₂ atmosphere in our laboratory (6).

cGMP/PDE activity assay in cell extracts. The cells were seeded at 1 \times 10⁶ cells/25-cm² flask. After 3 days, the cells were washed twice with phosphate-buffered saline (PBS), harvested with a rubber policeman, and homogenized in ice-cold homogenization buffer (1 mL) with T230 cell-freezing homogenizer (Tomy Seikagaku, Tokyo, Japan).

Dr. Piazza 2000 Patent



Date of Patent: *Dec. 5, 2000

Inventors: Rifat Pamukcu, Spring House; Gary A. Piazza, Doylestown, both of Pa.

activity of a test compound. Because the inventors have discovered a relationship between inhibition of cancer and inhibition of phosphodiesterase Type-5 isoenzyme (“PDE5”), this invention includes determining the PDE5

As used herein, the term “carcinoma” or “cancer” refers to lesions which are cancerous. Examples include malignant melanomas, breast cancer, prostate cancer and colon cancer.

Dr. Piazza 2018 Patent

Case 3:16-md-02691-RS Document 924-2 Filed 11/13/19 Page 5 of 94

(12) **United States Patent** (10) Patent No.: US 10,039,764 B2
Piazza (45) Date of Patent: Aug. 7, 2018

(54) TREATMENT AND DISEASES OR DISORDERS

(71) Application

(72) Inventor:

(73) Assign

(*) Notice:

(21) Appl. No.:	14/904,632	8,017,604 B2	9/2011	Albertari et al.
(22) PCT Filed:	Jul. 11, 2014	8,053,438 B2	11/2011	Allen et al.
(86) PCT No.:	PCT/US2014/044623	8,071,595 B2	12/2011	Ripka et al.

Examples of cancers characterized by solid tumors which may be treated include ...
melanoma

For instance, in some embodiments, the PDE10A inhibitor is co-administered with a PDE5 inhibitor to improve efficacy or reduce the effective dose range of the PDE10A inhibitory.

Representative examples of PDE5 inhibitors that may be useful in the practice of the present invention include sildenafil, tadalafil, vardenafil, udenafil, and avanafil or others such as MY5445 or compounds that increase intracellular cGMP levels (e.g., nitric oxide donors or releasing drugs). Examples of yet other PDE5 inhibitors that may be suitable for

(2013-01); A016 33/27 (2013-01); A016 45/06 (2013-01); C12Q 1/6886 (2013-01); G01N 33/5746 (2013-01); C12Q 2600/118

(62)

(3) ABSTRACT
Disclosed are methods for treating cancer and precancerous conditions with PDE10A specific inhibitors and diagnosis of cancer.

(58) Field of Classification Search

CPC A61K 31/415; A61K 31/4709; A61K 31/4745; A61K 31/4155; A61K 31/4409
See application file for complete search history.

8 Claims, 15 Drawing Sheets

Shawn P. Foley

(57) **ABSTRACT**
Disclosed are methods for treating cancer and precancerous conditions with PDE10A specific inhibitors and diagnosis of

CPC A61K 31/415; A61K 31/4709; A61K 31/4745; A61K 31/4155; A61K 31/4409
See application file for complete search history.

8 Claims, 15 Drawing Sheets

Ongoing Research

Nivolumab (Anti-PD1), Tadalafil and Oral Vancomycin in People With Refractory Primary Hepatocellular Carcinoma or Liver Dominant Metastatic Cancer From Pancreatic Cancers

The safety and scientific validity of this study is the responsibility of the study sponsor and investigators. Listing a study does not mean it has been evaluated by the U.S. Federal Government. Know the risks and potential benefits of clinical studies and talk to your health care provider before participating. Read our disclaimer for details.

Sponsor:
National Cancer Institute (NCI)

Information provided by (Responsible Party):
National Institutes of Health Clinical Center (CC) (National Cancer Institute (NCI))

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Brief Summary:

[https://clinicaltrials.gov/ct2/show/NCT03785210 \[9/9/2019 11:31:56 AM\]](https://clinicaltrials.gov/ct2/show/NCT03785210)

Sponsor: National Cancer Institute (NCI)

Tadalafil is a phosphodiesterase type 5 (PDE5) inhibitor . . .

PDE5 inhibitors have been examined in multiple malignancies and cancer cell lines for their **direct anticancer activities**, for their efficacy as **chemo-sensitizers** and for **cancer chemoprevention**.

PDE-5 Inhibitors and Immunity: Anti-Melanoma Effects

Chronic inflammation promotes myeloid-derived suppressor cell activation blocking antitumor immunity in transgenic mouse melanoma model

Christiane Meyer^{a,1}, Alexandra Sevko^{a,1}, Marcel Ramacher^{a,1}, Alexandre V. Bazhin^{a,b}, Christine S. Falk^{c,d}, Wolfram Oesen^a, Ivan Borello^a, Masashi Kato^d, Dirk Schadendorf^d, Michal Baniash^b, and Viktor Umansky^{a,2}

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“manipulation of the melanoma microenvironment with ...
sildenafil”

“significantly **increased survival** of tumor-bearing mice.”

ONCOIMMUNOLOGY
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<https://doi.org/10.1089/onco.2017.1326440>

ORIGINAL RESEARCH

Tadalafil has biologic activity in human melanoma. Results of a pilot trial with Tadalafil in patients with metastatic Melanoma (TaMe)



OPEN ACCESS



“**Tadalafil** led to a stabilization of the disease in 3 of 12 pretreated patients in the palliative setting.”

“Our study suggests that the PDE-5 inhibitor **tadalafil can improve clinical outcome of advanced melanoma patients. . .”**

immunogenicity of melanoma (3–6). However, despite the initial promising observations, the overall results of clinical immunotherapy trials are not satisfactory (3,5). Insufficient antitumor reactivity is thought to result from the formation of a complex immunosuppressive network induced by the chronic inflammation developing in the tumor microenvironment (6–8). Indeed, chronic inflammation has been demonstrated to correlate with tumor onset and progression (6, 7). An inflammatory microenvironment ensues during tumor growth as a result of the secretion of inflammatory mediators (cytokines, chemokines, growth factors, reactive oxygen and nitrogen species, prostaglandins) by the tumor and/or stroma cells (8–10). These mediators were found to support tumor development by stimulating protumorigen mutations, resistance to apo-

activity, and melanoma progression in *nr* transgenic mice. We found that an accumulation of functionally active MDSC in mel-

Author contributions: CM, AS, M.B., and V.U. designed research; CM, AS, M.R., A.V.L., A.V.S., and C.S.F. performed research; CM, AS, M.R., W.O., I.B., M.K., D.S., M.B., and V.U. analyzed data; and C.M., M.B., and V.U. wrote the paper.

The authors declare no conflict of interest.

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CM, AS, and M.R. contributed equally to this work.

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Supplementary information to this article can be accessed on the publisher's website.

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